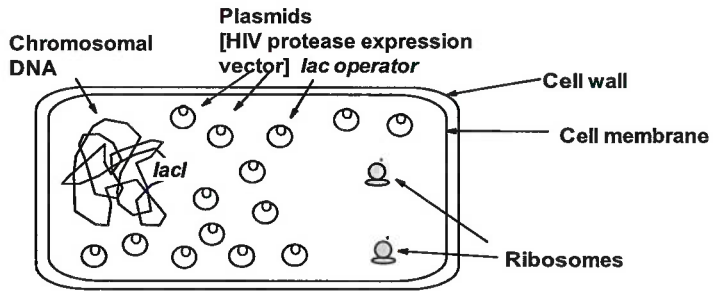


Lecture 39: Lac and T7 Expression Vectors

Lac Expression Vectors - Inducible

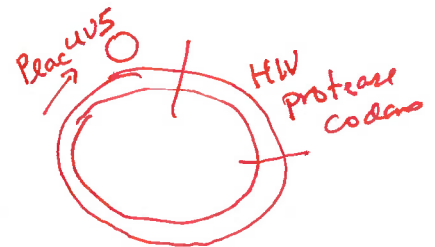
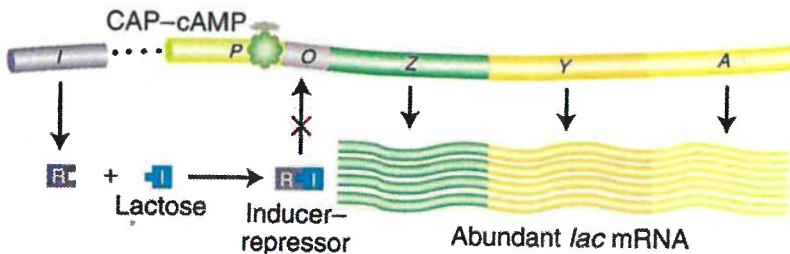
Expression of Recombinant Proteins:

- **Lac operator** – DNA sequence that binds the lac repressor protein, utilized as an on/off switch in the control of recombinant proteins.
- **Lac repressor** – protein that binds to the lac operator, released from DNA by IPTG
- **Lac I gene** – chromosomal, produces the lac repressor protein
- **LacUV5 promoter** – Mutation of the lac promoter that increases RNAP activity, avoiding the need for CAP+cAMP binding to get high levels of transcription.



1. The constitutive expression of high levels of almost any protein is toxic to the bacteria. The protein itself may be toxic, or the simple competition for cellular resources can lead to poor growth of the bacterial host, and in some cases, cell death. Therefore, it is necessary to regulate the production of high levels of recombinant protein.
2. In the Lactose operon production of proteins involved in the metabolism of lactose are controlled by the binding of the **lac repressor** (a protein that is the product of the *lacI* gene, produced from the bacterial chromosome) to a region of the DNA near the promoter region of the genes that encode the proteins for lactose metabolism. This segment of DNA is called the **lac operator**. High-level expression from the lac operon only occurs when glucose is low, leading to high cAMP levels in the bacterial cytoplasm. cAMP binds to the catabolite activator protein (CAP), the CAP-cAMP complex binds upstream from the lac promoter enhancing transcription.

(c) No glucose present (cAMP high); lactose present



3. The lac operator system can be used it to control expression of any other gene by simply placing the appropriate DNA segments in the correct location in our expression vector. To avoid regulation by CAP, the normal lac promoter is replaced by the lacUV5 promoter, this gives high levels of expression even if glucose levels are high.

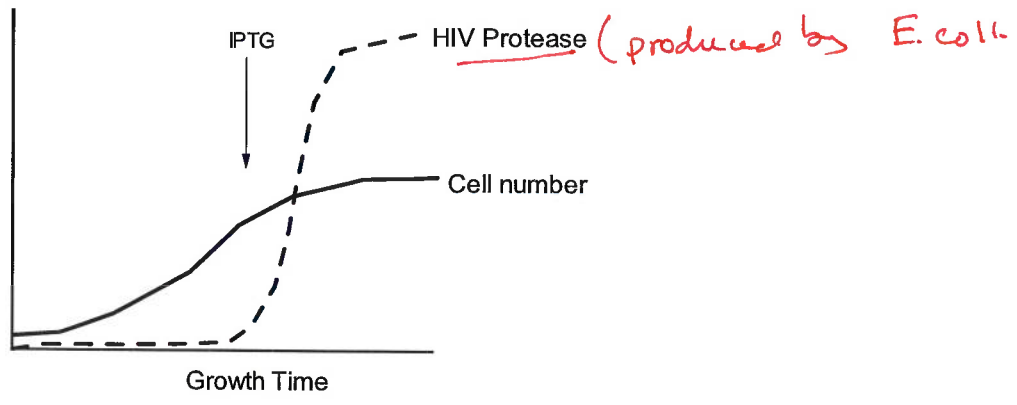
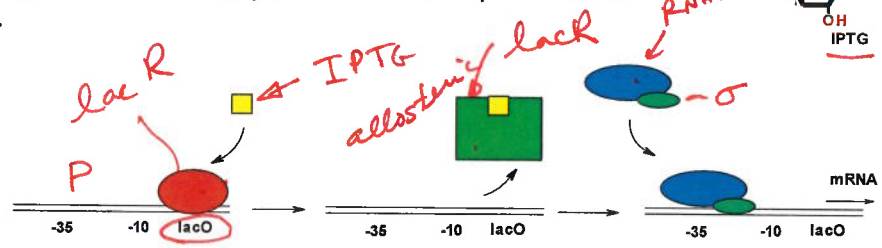
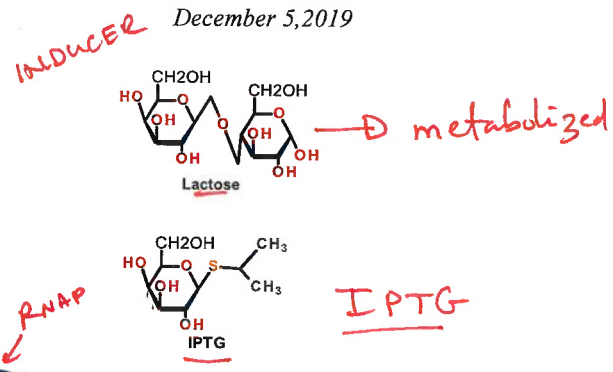
```

AATGTGAGTTAGCTCACTCATTAGGCACCCAGGCTTTTACACTTTATGCTTCCGGCTCGTATGTTGTGTGTGGAATTGTGAGCGGATAACAATTAGGAAACAGCTATG...
CAP binding                               Lac promoter                               Protein->
AATGTAAGTTAGCTCACTCATTAGGCACCCAGGCTTTTACACTTTATGCTTCCGGCTCGTATAATGTTGTGTGGAATTGTGAGCGGATAACAATTAGGAAACAGCTATG...
-35                               -10                               mRNA----->
LacUV5 promoter                   <--lac operator----->                               Met...
    
```

Reflection: why is it preferable to use the lacUV5 promoter instead of the normal lac promoter?

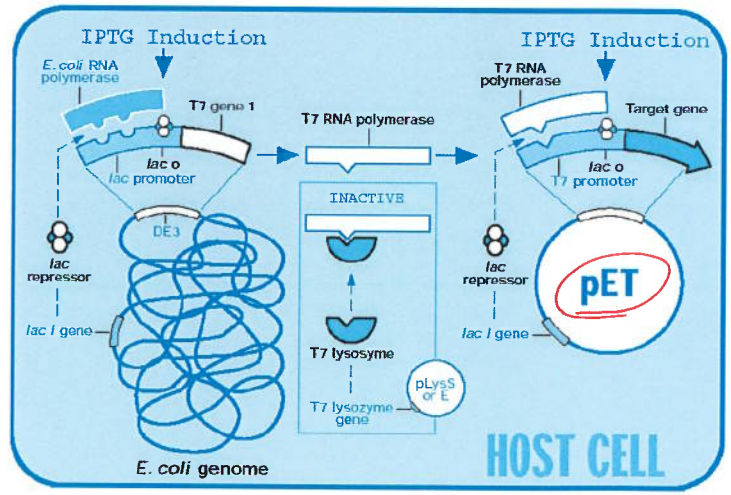
use high glucose → optimal growth.

4. The lac repressor binds to the DNA when lactose is absent and blocks transcription of the DNA. When lactose is present, it binds to the lac repressor, causing an **allosteric** change that releases it from the DNA. Since lactose would be rapidly degraded by the bacteria, a non-hydrolyzable analog, isopropylthio-galactoside (IPTG), is used instead. Once the lac repressor leaves the DNA, RNA polymerase can bind, allowing production of mRNA that can be used by the ribosome to produce HIV protease.

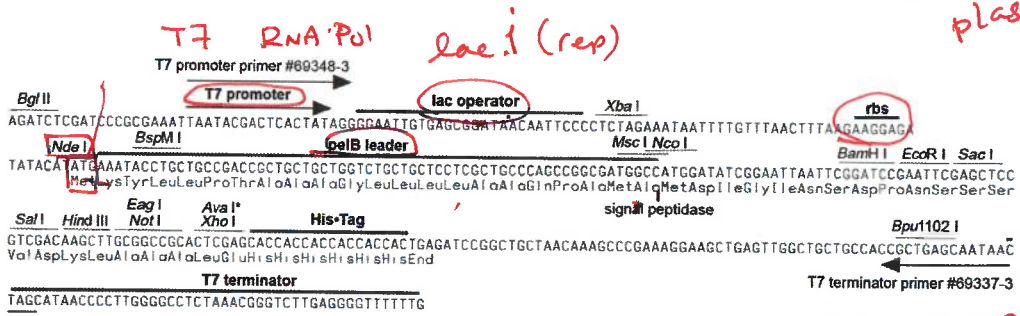
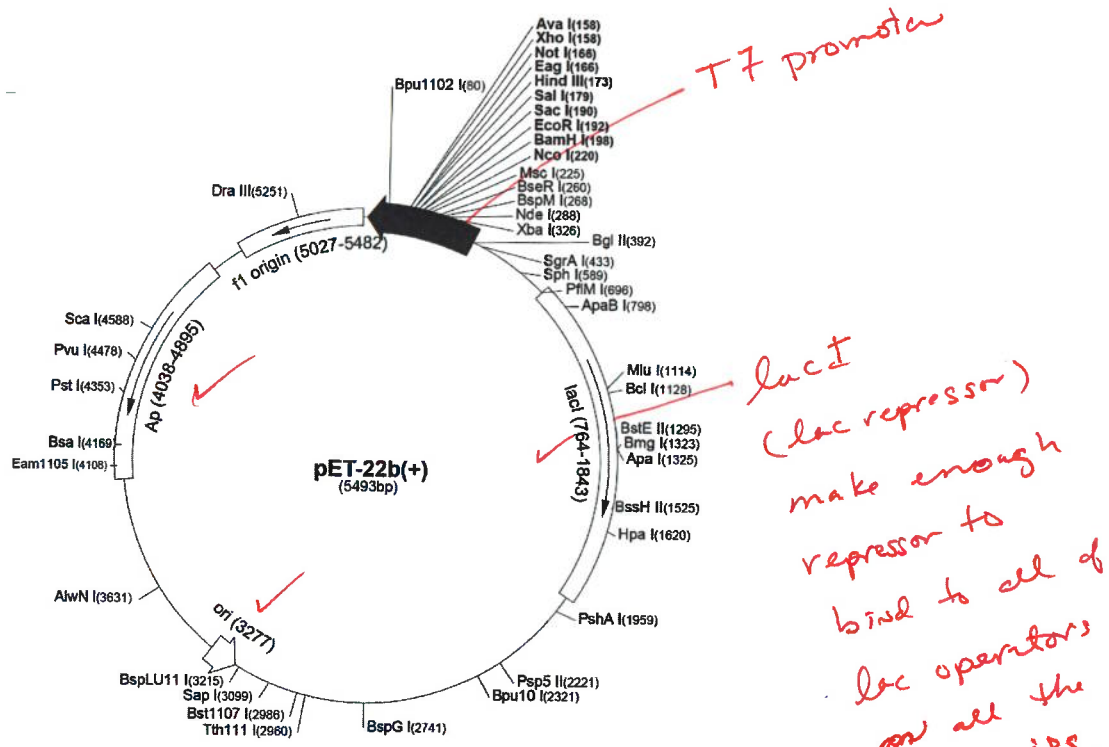


Alternative Expression Systems: T7-pET vector Series.

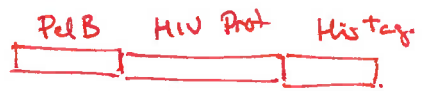
- Use T7 phage promoter/polymerase. Only significant difference is a different promoter is used to drive production of mRNA and a specialized host strain is required that expresses T7 polymerase.
- T7 polymerase only recognizes T7 promoters, and is more active than *E. coli* RNA polymerase, producing more mRNA transcripts (and therefore more protein).
- T7 polymerase under lac control on the host *E. coli* chromosome. T7 polymerase only produced when IPTG is added. 1st level of control in expression.
- Target gene is usually also under lac control on the plasmid. 2nd level of control of expression.
- Production can be attenuated by T7 lysozyme gene which inactivates T7 RNA polymerase.



- i) IPTG
- ii) produce T7 RNA Pol
- iii) produce mRNA target gene (HIV protease)



pET-22b(+) cloning/expression region



1. Expression from the T7 promoter is under double lac control, the lac repressor blocks expression of T7 polymerase (on the chromosome) and expression of the gene on the plasmid.
 2. Inserted codons using the NcoI site will add a signal peptide to the protein to cause expression from the cell.
 3. A His-tag can also be added to the C-terminus of the protein, using the stop codon after the six His residues.
- Cloning into pET22b using NcoI/XhoI would use both of these features. The PCR product that would be inserted would have the following sequence.

YYCCATGGXX[codon 1][codon 2].....CTCGAGZZ
 YYGGTACC GAGCTCZZ
 NcoI Xho I

- The extra YY and ZZ bases allow NcoI and Xho I to efficiently cleave the DNA (many restriction enzymes cannot cleave if their site is at the very end of a fragment).
- The three bases (GXX) after the underlined ATG represent the 2nd codon, which is restricted to Val, Ala, Asp, Glu, Gly in order to have the NcoI site.

The final construct will be: -T7-lacO-rbs-Met---NcoI-coding region-Xho-HisHisHisHisHisHisTAA-----

PelB - Export of Recombinant Proteins.

A specialized *protein* sequence, called the leader peptide, when present as the amino terminal residues of the protein, signals the export of the protein out of the cell. This may reduce the toxicity of the protein and make purification easier since only a small number of proteins are exported out of the cell.

During the export process, this peptide is cleaved by the **leader peptidase** (also known as signal peptidase), producing the mature exported peptide.

The main features of this peptide are:

- i) Basic residue at amino terminus
- ii) Non-polar segment of ~15 amino acids.
- iii) Cleavage site, which is followed by a second basic residue (---Ala[^]Arg---

To use pET22b to express HIV protease we would modify our original PCR primers to add the NcoI and XhoI sites (instead of the EcoR1 and BamH1), after ligating into pET the final sequence, from the ribosome binding site to the stop codon is:

```

RBS      NdeI      NcoI      XhoI
AAGGAGATATACATATGAAATACCTGCTGCCGACCGCTGCTGGTCTGCTGCTCCTCGTCCAGCCGCGCATGGCATGGAT--HIV Protease---CTCGAGCACCACCACCACCACCTGA
MetLysTyrLeuLeuProThrAlaAlaAlaGlyLeuLeuLeuLeuAlaAlaGlnProAlaMetAlaMetAspProGln-----AsnPheLeuGluHisHisHisHisHisHis
    
```

2) The peptide, as it comes off of the ribosome will look like:

```

-----Leader Sequence-----|-----HIV protease-----| |.....His Tag....
fMetLysTyrLeuLeuProThrAlaAlaAlaGlyLeuLeuLeuLeuAlaAlaGlnProAlaMetAlaMetAspProGln-----AsnPheLeuGluHisHisHisHisHisHis
                                     Cleavage site for leader peptidase
    
```

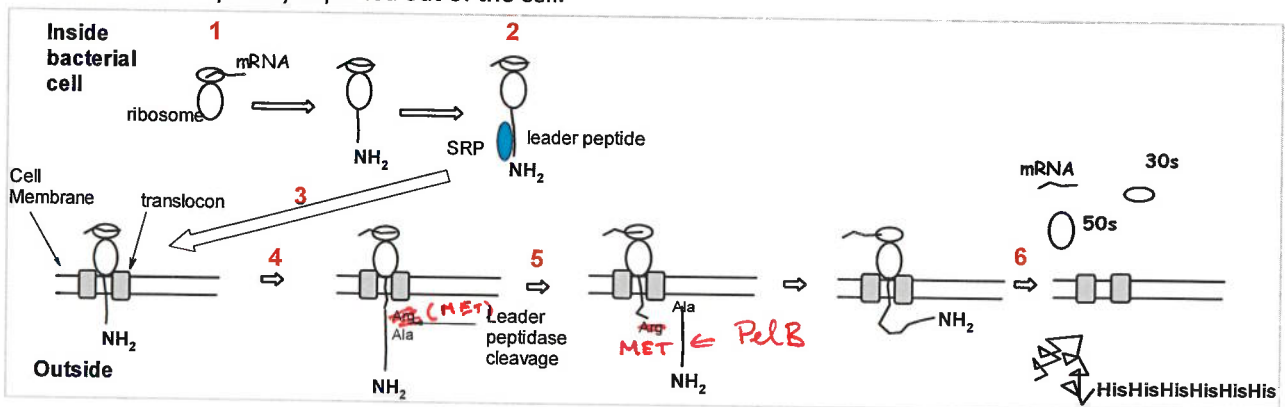
3) After export out of the cell the final product will be as follows (The bold amino acids were added as part of the NcoI and XhoI sites that were required to insert the PCR product into the expression vector.):

```

|-----HIV protease-----| |-----His Tag-----|
MetAspProGlnIle----LeuAsnPheLeuGluHisHisHisHisHisHis
    
```

Protein Export Machinery:

1. mRNA binds to the ribosome.
2. When leader peptide emerges from the ribosome it binds to the signal recognition particle (SRP).
3. The ribosome/mRNA/SRP binds to transport machinery(translocon) in the cell membrane.
4. Protein synthesis continues, protein extruded outside the cell.
5. Leader peptidase cleaves off leader peptide.
6. Protein is completely exported out of the cell.



Note: The cleavage site for the peptidase is either Ala-Met, or Ala-Arg, Ala-Met is found in the PelB leader.

